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Name (Print)

Signature

Customer No.: 07278

Docket No: 03394/100H557-US1

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: Ehud Goldin et al.

Serial No.:

09/851,494

Art Unit:

1646

Confirmation No.:

Filed:

For:

Examiner:

John D. Ulm

Gene Encoding A New TRP Channel Is Mutated In Mucolipodosis IV

DECLARATION UNDER 37 C.F.R. § 1.131

Mail Stop Non-Fee Amendments Commissioner for Patents P.O. Box 1450 Alexandria, VA 22313-1450

Sir:

- I, Susan A. Slaugenhaupt, hereby declare and state as follows:
- I am a citizen of the United States of America. I am more than twenty-

one years of age.

Serial No. 09/851,494

Docket No: 03394/100H557-US1

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- 2. I am a co-inventor of the above-identified application, along with James S. Acierno JR. James S. Acierno JR and I did the work that resulted in the data reported in the declaration, or it was done under my supervision as head of the laboratory.
- 3. I make this statement on behalf of myself and the co-inventors identified in paragraph 2.
- 4. I reaffirm my duty of candor and good faith in dealing with the Office, including the duty to disclose to the Office all information known to be material to the patentability of the invention as defined in 37 C.F.R. § 1.56.
- 5. I have read and am familiar with the instant application as it was filed in the U.S. Patent and Trademark Office.
- 6. I have read and am familiar with the publications by (i) Curtis et al. (Pub. No. US 2002/0035056 A1), which has an effective filing date under 35 U.S.C. 119(e) of Apr. 07, 2000; and (ii) Lal et al. (Pub. No. US 2002/0182671 A1), which has an effective filing date under 35 U.S.C. 119(e) of Aug. 17, 1999.
- 7. It is my understanding that, according to the Examiner, the amino acid sequence presented in SEQ ID NO: 3 of the instant application is identical to the amino acid sequence presented in SEQ ID NO: 2 of Curtis et al. and SEQ ID NO: 13 of Lal et al. It is further my understanding, that the Examiner states that Curtis et al. and Lal et al. each present an isolated nucleic acid encoding a protein comprising the amino acid sequence presented in

Serial No. 09/851,494

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SEQ ID NO: 3 of the instant application, as well as a vector and host cell comprising that nucleic acid.

- 8. Prior to Aug. 17, 1999, the effective date of the Lal et al. publication, we conceived and reduced to practice the invention as described and claimed in claims 1, 5-7, and 33-34 of the subject application.
- 9. As evidence that our reduction to practice antedates Lal et al., we refer to Exhibits 1 and 2, which collectively establish the conception and reduction to practice of our invention prior to Aug 17, 1999. The exhibits document isolation and possession of a nucleic acid encoding MCOLN1 prior to Aug. 17, 1999. Dates, along with privileged information, appearing in these documents have been reducted, but each document has a date before August 17, 1999.
- 10. Exhibit 1 establishes identification of MCOLN1 sequence, showing the receipt by Dr. Slaugenhaupt of two primers: (i) sts-T66288-R (5'-AGC TGC AGG CCT ACA TCG -3'); and (ii) sts-T66288-F (5'GGC AGT CAG GTC GAA TCA AT-3). As shown in Appendix A, the two primers are specific to the MCOLN1 gene, spanning the 1732-1883 bp region of the MCOLN1 cDNA sequence (SEQ ID NO: 3).
- 11. Exhibit 2 documents identification and possession of a nucleic acid encoding a full-length MCOLN1 protein by showing an EST alignment spanning the MCOLN1 gene. At least two notations are particularly relevant. First, this page shows a "2264 bp"

Serial No. 09/851,494

Docket No: 03394/100H557-US1

annotation of T66288 following sequencing, indicating that T66288 encodes the entire MCOLN protein. Prior to our sequencing, the exact insert size of this construct was not known. Second, this page also identifies the orientation of AI8166064, which is the corresponding GenBank accession number for IMAGE CLONE 2517653 (Appendix B). Paragraph [0185] of the specification states that we "sequenced the IMAGE clone 2517653." This paragraph further describes our deduction and confirmation of the MG-2 (MCOLN) open-reading frame from this clone.

- also achieved reduction to practice of an expression vector encoding the MCOLN1 protein prior to August 17, 1999. Appendix B reveals that IMAGE CLONE 2517653 (as presented in Exhibit 2) is inserted into the pBlusescript SK+ vector. This common vector is widely recognized by those skilled in the art of molecular biology as including T3 and T7 promoters that flank the cloning site, which allow expression of the inserted gene sequence. Appendix C shows the key structural features of this vector. The entire MCOLN1 open reading frame is present in IMAGE CLONE 2517653. Alternatively, given our identification of the coding sequence of MCOLN1, we knew how to incorporate that sequence into an expression vector such as pBluescript SK+.
- 13. These documents establish that the inventions of claims 1, 5-7, and 33-34 were reduced to practice prior to Aug. 17, 1999.
- 14. I further declare that all statements made herein of my own knowledge are true, and that all statements made on information and belief are believed to be true. I

 Serial No. 09/851,494

 Docket No: 03394/100H557-US1
 Page 4

further declare that these statements are made with the knowledge that the willful false statements and the like so made are punishable by fine or imprisonment or both, under Section 1001 of Title 18 of the United Stated Code, and that such willful false statements may jeopardize the validity of the instant application or of any patent issued thereupon.

Respectfully submitted,

DATE

Susan A. Slaugenhaupt

Serial No. 09/851,494

Docket No: 03394/100H557-US1

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PAGE 1 EXHIBITI

grated DNA Technologies, Inc.

gonucleotide Specification Sheet

stomer Information

Susan Slaugenhaupt Harvard Institute of Human Genetics Massachusetts General Hospital-Boston 77 Avenue Louis Pasteur HIM Bldg. Rm. 422 Boston, MA 02115 6174327025

Mail: order //www.idtdna.com

Order Information

Order Date:

Customer #:

19479

P.O. #:

0000085288

Sales order #: 148396 624757 Reference #:

Oligonucleotide Information

Reference #: Purification:

624757

Standard Purification

Sequence Name: sts-T66288-f

Product: DNA Oligo Sample

Unit Size: 100 nmole

Bases: .20

5'- GGC AGT CAG GTC GAA TCA AT -3" Sequence:

Molecular Weight:

7,572.00

GC Content:

50.0 %

Tm (50mM NaCl):

51.44 °C

Amount of Oligo

21.8

95.01

0.72

OD260

nanomoles

ma

Printed

6/9/99



624757 Invegrand DNA Tech 6. Slaugenhoupt os/08/89 .

BB2BB4 C AGT CAG GTC CAA TCAAT Tm = 51.44 °C, MW = 7572 21.80 00= - 95.01 cmd + 0.72 cm

624757 Interest DNA Tech 100/09/99 OS/09/99 S. Shrippenhark odjus/99 mp-tecses/ B-dod act ord uto was tea at 4 Tri = 5154 -0, MY = 7572 21.80 Oca = 95,01, maiol = 0.72 mg

Samples Statistically Tested

Q.C. Approved By:

PLEASE READ BEFORE OPENING TUBES

- Store at -20°C. If the oligo is to be used on multiple occasions, resuspend in water or tris-EDTA buffer, divide into smaller aliquots, lyophilize, and store at -20°C.
- Contents may appear as either a translucent film or a white powder. This variance does not affect the quality of the oligo. * Please centrifuge tubes prior to opening. Some of the product may have been dislodged during shipping.
- Calculations are made using 1 $\overrightarrow{OD}_{200} = 33 \text{ ug} / \text{mL}$

/www.ldtdna.com

igonucleotide Specification Sheet

istomer information

Susan Slaugenhaupt Harvard Institute of Human Genetics Massachusetts General Hospital-Boston 77 Avenue Louis Pasteur HIM Bldg. Rm. 422 Boston, MA 02115 6174327025

Order Information

Order Date: Customer #:

P.O. #:

19479

0000085288

Sales order #: 148396 Reference #: 624758

Oligonucleotide information

Reference #:

624758

Purification:

Standard Purification

Sequence Name: sts-T66288-R

Product: DNA Oligo Sample

Unit Size: 100 nmole

Bases:

18

Sequence: 5'- AGC TGC AGG GCT ACA TCG -3'

Molecular Weight:

6,754.00

GC Content:

61.1 %

Tm (50mM NaCl):

51.11 °C

Amount of Oligo

15.5

75.73

0.51

OD₂₆₀

nanomoles

mg

Printed 6/9/99

LABELS - PEEL HERE

624758 Integrated DNA Tech S. Shauperticupt 08/08/68 ma-186285-R EACC FOR ASS COT ACA TOO A in +31.11 °C, MW - 6754

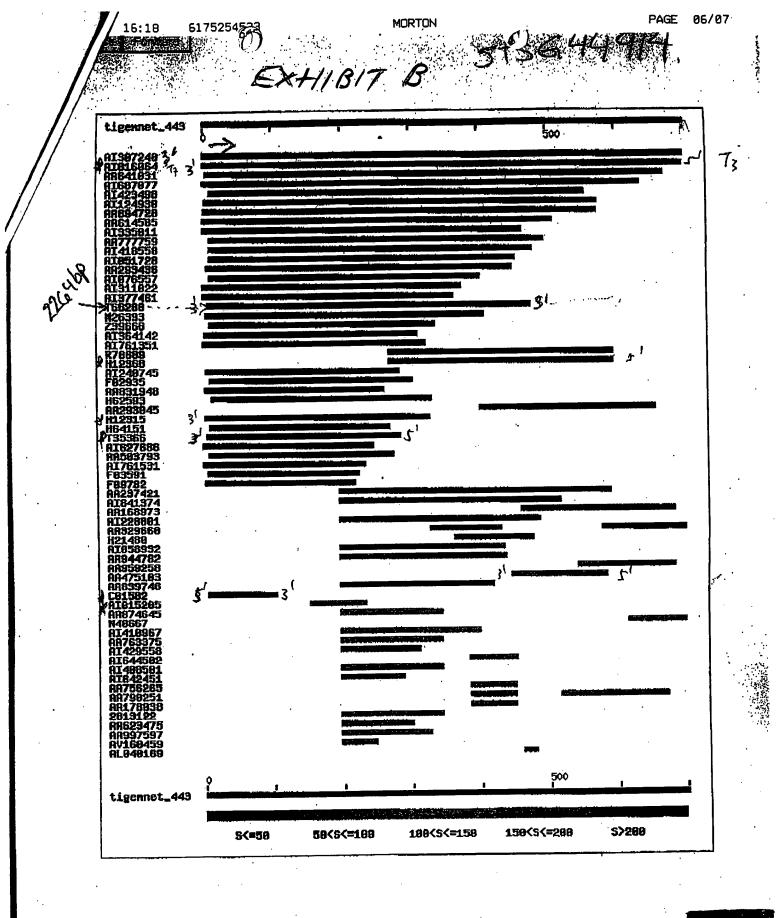
624758 Integrated DNA Tech 8. Skrugenhaupt 09/09/95 89-768288-R 8'-AGC TGC AGG GCT AGA TGC 48

Samples Statistically Tested Q.C. Approved By:

PLEASE READ BEFORE OPENING TUBES

Store at -20°C. If the oligo is to be used on multiple occasions, resuspend in water or tris-RDTA buffer, divide into smaller aliquots, lyophilize, and store at -20°C.

Contents may appear as either a translucent film or a white powder. This variance does not affect the quality of the oligo. Please centrifuge tubes prior to opening. Some of the product may have been dislodged during shipping. Calculations are made using 1 OD 26 = 33 ug / ml



Page 1 APPENDIX A

SEQ ID NO: 2

------ sts-T66288-f ------ sts-T66288-r <400>

960	gagacccagg	catcagcctg	ggcggatccc	gcacacagtg	tgacaacaaa	tgatcacgtt
006	ttcagcgtcc	ctgctatacc	agatcccgga	atcaataatg	ccagagcctc	ccattaacct
840	cggctgaaga	catccacttc	tcaatgtcac	cacaagctgg	gctcaaattc	agaacctcac
780	tccagttaca	ggaaagcagc	tcaccctctt	agcgacgatc	tccgccccc	адсддсссс
720	gatccccccg	catccaggtg	ttactgactg	ccgatggtgg	tgacattgat	acgacacatt
099	gacccggcca	aggccacgtg	actaccaccg	tgccagcggt	gcttgctctc	atggctcagg
009	ccttggacca	tgggggtgac	atgtccgtgg	cggtatgcgt	gtcactgggc	tgcctgacgt
540	tacctggcgt	tgtggaccag	tcttccatgc	taccaggcca	ggagcagctg	cctacacgcg
480	accttcgcag	agcggatgac	actcggacgg	acacctcttc ctgctgggct		tegeetteeg
420	gagaacacca	attccgggaa	tggctgtgac	agtaatcagc	gtttgggctc	agctcatcct
360	gtcacggtgc	gatcctggtg	aagtggtcaa	ctgatgctgc	gccctgcaag	адддссдсаа
300	tttcgagcca	ctgcgacaag	tcatgagtcc	aaatactttt	ccgtcgtctc	aagaccttcg
240	ccagaagagg	tccgacaccc	caccggcccc	gcggggcctt	tgggacccag	accccgggta
180	ctgaccccca	cgagcggctt	gctcagagac	ggtccgcgcg	agccccggcg	ccagcatgac
120	cgctcccgcc	ctgcgtcccg	ccgtacccgc	ggctgccccg	atcggaccca	მამნმანმან
09	cgcagtgaca	ggagcgaggt	gcgccgcgag	gtttgaagcc	atgccggagg	agatcagctg



Page 2 APPENDIX A

J. 13 . L

2051					ಹ	tgttgaataa
204(gtggggaggg	gggagactgg	cttgggggcg	gtgtcggacc	tggacctttc	gaggaggcc
198(tgtcgcgccc	atcggctccc	gcttttaagg	tgtagggttt	cttatttatt	cccgaccccg
1920	gacccccgcc	ggactgcaga	cgtaggccct	gccgttggac	cgacctgact	tgaattgatt
1860	tcgctgctgg	ggaggagcat	gggacccctc	tgctgcggaa	ccttctctgc	cggcctgcag
1800	gggagcggct	gttccgccgc	cctccggcaa	gacagcccca	acagtgccag	cctacatego
174(gagctgcagg	agaggagagc	gcgcaggcgc	catcccggcg	caccatcaag	gcgcctacga
1680	ctcatcaccg	cttcatcgcg	tgctcagcct	atctacatgg	cagcctcttc	actccttcat
1620	ctctaccttt	cttctcccag	tggtgtggct	cgcagcagcc	gcagcagggc	ccatgcaggc
1560	acgttcgccg	catgtttgtg	atggggacga	tegeteatea	gtgcctgttc	tggtgtctga
1500	tcactctcca	gaagttccgc	cctatcatgt	gtgctggggc	tggctggatc	actgcttctg
1440	tacctgggct	ggctgtcatc	gctgctgcgt	atgcgcttct	gcccagcgtc	gggtggccct
1380	gccacactgc	tatcctcatc	acaactacaa	accttcttcc	ccgctacctg	tgggcgtgat
1320	ctggtgtggg	ctcgacgctg	tcctgggcac	tgcagcatcc	ctacgacgtc	acttggcgag
1260	gaggccaaga	gatcggcatc	ccatcatgaa	atctcgggca	tgtgctcacc	tcaccagcga
1200	atcctgctcg	tggctggtac	aatttgtcaa	gagcggctgg	cagcctgtgg	gacgggtcat
1140	cggcagcggg	gttcatgtgg	agtttgtggg	ctgcagaacg	aggcttcctg	cactccttcg
1080	tgagaaagat	cttcctcctc	gctccctgtc	atcctcacct	cgtggtggtc	tcctgtttga
1020	agcttccggc	cggagacaac	tcttccagca	caccccagtg	ggagtgtaag	cccacatcca

 $\left\{W.: 03394 \setminus 100H557US1 \setminus 00204873. DOC \text{ IN MINIMUM IN MINIMUM IN MINIMUM }\right\}$

I.M.A.G.E. Clone Query: Results



Blotechnology & Shorechnology A. B. Consortium of R. A. B. Consortium

"Sharing resources to achieve a common goal - the discovery of all genes "

Begin new search | Begin new Clone search

I.M.A.G.E. Clone Query Results:

Your search returned 2 results! Here they are:

Result Number	CLONE	ROW POS	COL	PLATE	GB ACCNUM	SEQ LENGTH	GB DATE CREATED	$\begin{array}{c c c c c c c c c c c c c c c c c c c $	CDNA LIBR ID	CDNA LIBR SPECIES ID	TISSUE	VECTOR NAME
1	2517653	I	9	6268 AI81	AI815981	448	Jul 09 1999 12:00AM	Jul 09 1999 Apr 17 2003 12:00AM 05:06PM	1341	human	brain/CNS	brain/CNS pBluescript SK+
2	2517653	1	9	6268 AI81	AI816064	902	Jul 09 1999 12:00AM	Jul 09 1999 Apr 17 2003 12:00AM 05:06PM	1341	human	brain/CNS	brain/CNS pBluescript SK+

Begin new search | Begin new Clone search

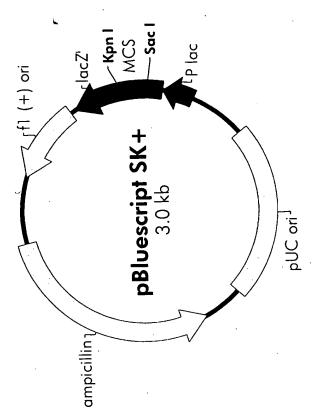
I.M.A.G.E. Consortium home page

BBRP home page

LLNL Programs, Projects, Centers and Consortia

APPENDIX (

f1 (+) origin 138–444 β -galactosidase α -fragment 463–816 multiple cloning site 653–760 lac promoter 817–938 pUC origin 1458–1825 ampicillin resistance (bla) ORF 1976–2833



pBluescript SK (+/-) Multiple Cloning Site Region (sequence shown 601–826)

. ATCGATAAGCTTGATATCGAATTCCTGCAGCCCGGGGGATCCACTAGTTCTAGAGCGGCCGCCACCGCGGTGGAGCTCCA.... SK primer binding sile TTGTAAAACGACGGCCAGTGAATTGTAATACGACTCACTATAGGGCGAATTGGGTACCGGGCCCCCCCTCGAGGTCGACGGT.

KS primer binding sile

KS primer binding sile | Fag | BstX | Sac || Sac | Hinc II Sol I Xhol Apo I EcoO109 I Ora II Kpn I gamH1 Spe1 Xba1 T7 Promoter € Smal Bsp106 | Hind III EcoRV EcoRI Pst |

β-gal α-fragment

GCTTTTGTTCCCTTTAGTGAGGGTTAATTTCGAGCTTGGCGTAATCATGGTCATAGCTGTTTCC

M13 Reverse primer binding site